



Kuroda Revised Sequence Listing filed 2006-08-14  
SEQUENCE LISTING

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Kosugi, Yoshinori  
Ohbayashi, Tetsuya

<120> Methods for Diagnosing Endometriosis-Related Diseases

<130> 2006\_0025A

<140> 10/564,481

<141> 2006-01-13

<150> JP 2003-196455

<151> 2003-07-14

<160> 8

<170> PatentIn version 3.3

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<213> Homo sapiens

<220>

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Met Ile Ile Tyr Arg Asp Leu  
1 5

atc agc cac gat gag atg ttc tcc gac atc tac aag atc cgg gag atc 163  
Ile Ser His Asp Glu Met Phe Ser Asp Ile Tyr Lys Ile Arg Glu Ile  
10 15 20

gcg gac ggg ttg tgc ctg gag gtg gag ggg aag atg gtc agt agg aca 211  
Ala Asp Gly Leu Cys Leu Glu Val Glu Gly Lys Met Val Ser Arg Thr  
25 30 35

gaa ggt aac att gat gac tcg ctc att ggt gga aat gcc tcc gct gaa 259  
Glu Gly Asn Ile Asp Asp Ser Leu Ile Gly Gly Asn Ala Ser Ala Glu  
40 45 50 55

ggc ccc gag ggc gaa ggt acc gaa agc aca gta atc act ggt gtc gat 307  
Gly Pro Glu Gly Glu Gly Thr Glu Ser Thr Val Ile Thr Gly Val Asp  
60 65 70

att gtc atg aac cat cac ctg cag gaa aca agt ttc aca aaa gaa gcc 355  
Ile Val Met Asn His His Leu Gln Glu Thr Ser Phe Thr Lys Glu Ala  
75 80 85

tac aag aag tac atc aaa gat tac atg aaa tca atc aaa ggg aaa ctt 403  
Tyr Lys Lys Tyr Ile Lys Asp Tyr Met Lys Ser Ile Lys Gly Lys Leu  
90 95 100

gaa gaa cag aga cca gaa aga gta aaa cct ttt atg aca ggg gct gca 451  
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Glu Glu Gln Arg Pro Glu Arg Val Lys Pro Phe Met Thr Gly Ala Ala  
 105 110 115

gaa caa atc aag cac atc ctt gct aat ttc aaa aac tac cag ttc ttt 499  
 Glu Gln Ile Lys His Ile Leu Ala Asn Phe Lys Asn Tyr Gln Phe Phe  
 120 125 130 135

att ggt gaa aac atg aat cca gat ggc atg gtt gct cta ttg gac tac 547  
 Ile Gly Glu Asn Met Asn Pro Asp Gly Met Val Ala Leu Leu Asp Tyr  
 140 145 150

cgt gag gat ggt gtg acc cca tat atg att ttc ttt aag gat ggt tta 595  
 Arg Glu Asp Gly Val Thr Pro Tyr Met Ile Phe Phe Lys Asp Gly Leu  
 155 160 165

gaa atg gaa aaa tgt taa caaatgtggc aattattttg gatctatcac 643  
 Glu Met Glu Lys Cys  
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ctgtcatcat aactggcttc tgcttgtcat ccacacaaca ccaggactta agacaaatgg 703

gactgatgtc atcttgagct cttcatttat tttgactgtg atttatttgg agtggaggca 763

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Gly Lys Met Val Ser Arg Thr Glu Gly Asn Ile Asp Asp Ser Leu Ile  
 35 40 45

Gly Gly Asn Ala Ser Ala Glu Gly Pro Glu Gly Glu Gly Thr Glu Ser  
 50 55 60

Thr Val Ile Thr Gly Val Asp Ile Val Met Asn His His Leu Gln Glu  
 65 70 75 80

Thr Ser Phe Thr Lys Glu Ala Tyr Lys Lys Tyr Ile Lys Asp Tyr Met  
 85 90 95

Lys Ser Ile Lys Gly Lys Leu Glu Glu Gln Arg Pro Glu Arg Val Lys  
 100 105 110

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Pro Phe Met Thr Gly Ala Ala Glu Gln Ile Lys His Ile Leu Ala Asn  
115 120 125

Phe Lys Asn Tyr Gln Phe Phe Ile Gly Glu Asn Met Asn Pro Asp Gly  
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